Application No. 09/970,966

Amendment Dated September 3, 2003

Reply to Final Office Action dated May 5, 2003

## Amendments to the Claims:

This listing of claims will replace all prior versions, and listings, of claims in the application:

## Listing of Claims:

- 1-21. (Canceled)
- (Amended) A method for determining the presence of ovarian cancer in a patient, comprising the steps of:
- (a) contacting a biological sample obtained from a patient with a probe-consisting of at least 10 contiguous nucleutides of a sequence selected from the group consisting of:
  - a) ahout 20 to about 363 configuous nucleotides of SEQ ID

NO:199.

b) about 20 to about 1917 continuous nucleotides of SEQ ID

NO:214, and

- c) the complete complements of a) and b);
- (b) detecting in the sample an amount of an expressed polynucleotide that hybridizes to the probe under moderately stringent conditions; and
- (c) comparing the amount of expressed polynucleotide that hybridizes to the probe to a predetermined cut-off value, and therefrom determining the presence of ovarian cancer in the patient.
- 23. (Amended) A method for determining the presence of ovarian cancer in a patient, comprising the steps of:
- (a) contacting a biological sample obtained from a patient with at least two oligonucleotide primers, each primer comprising consisting of at least 10 to 363 contiguous nucleotides of SEQ ID NO:199 or the complete complement thereofor 10 to 363 contiguous nucleotides of SEQ ID NO:199, in a reverse transcriptase polymerase chain reaction, wherein

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said oligonucleotide primers are capable of amplifying a polynucleotide sequence recited in SEQ ID NO:199; and

- (b) detecting in the sample an amount of an expressed polynucleotide sequence that amplifies in the presence of said oligonucleotide primers;
- (c) comparing the amount of expressed polynucleotide that amplifies in the presence of said oligonucleotides to a pre-determined cut off value, and therefrom determining the presence of ovarian cancer in the patient.
- 24. (Amended) A method for determining the presence of ovarian cancer in a patient, comprising the steps of:
- (a) contacting a biological sample obtained from a patient with at least two oligonucleotide primers, each primer comprising at least consisting of 10 to 1917 contiguous nucleotides of SEQ ID NO:214 or the complete complements thereofof 10 to 1917 contiguous nucleotides of SEO ID NO:214, in a reverse transcriptase polymerase chain reaction, wherein said oligonucleotide primers are capable of amplifying an expressed polynucleotide sequence recited in SEQ ID NO:214; and
- (b) detecting in the sample an amount of an expressed polynucleotide sequence that amplifies in the presence of said oligonucleotide primers;
- (c) comparing the amount of expressed polynucleotide that amplifies in the presence of said oligonucleotides to a pre-determined cut off value, and therefrom determining the presence of ovarian cancer in the patient.
- 25. (New) The method of claim 22, wherein the probe is selected from the group consisting of a) 25 to 363 contiguous nucleotides of SEQ ID NO:199, b) 25 to 1917 contiguous nucleotides of SEQ ID NO:214, and c) complete complements of a) and b).
- 26. (New) The method of claim 32, wherein the probe is selected from the group consisting of a) 50 to 363 contiguous nucleotides of SEQ ID NO:199, b) 50 to 1917 contiguous nucleotides of SEQ ID NO:214, and c) complete complements of a) and b).